

(FILE 'MEDLINE, CANCERLIT, EMBASE, BIOSIS, BIOTECHDS, CAPLUS' ENTERED AT  
13:28:29 ON 01 AUG 2002)

DEL HIS

L1 72776 S ANTHRACENE  
L2 1648352 S MUTA? OR HYPERMUTA? OR MISMATCH REPAIR  
L3 5117 S L1 AND L2  
L4 13402586 S CELL# OR MAMMAL OR RAT OR MURINE OR MOUSE OR IN VIVO  
L5 3606 S L4 AND L3  
L6 1669 DUP REM L5 (1937 DUPLICATES REMOVED)  
L7 3791 S HYPERMUTAT?  
L8 0 S L7 AND L6  
L9 992579 S MUTAT?  
L10 2871 S L9 AND L7  
L11 12130 S MISMATCH REPAIR  
L12 144 S L11 AND L10  
L13 672 S L9 AND L6  
L14 2 S L13 AND L11  
L15 398 S L13 AND CELL  
L16 4984 S 1,2-DIMETHYL  
L17 47 S L16 AND L1  
L18 8 S L17 AND L4  
L19 6 DUP REM L18 (2 DUPLICATES REMOVED)  
L20 399 S L6 AND ASSAY  
L21 56 S L20 AND GENE

=>

L15 ANSWER 198 OF 398      CANCERLIT

AN 97600055      CANCERLIT

DN 97600055

TI P53 **mutations** in chemically induced hamster cheek-pouch tumors  
(Meeting abstract).

AU Gimenez-Conti I B; LaBate M; Liu F; Osterndorff E  
CS Dept. of Carcinogenesis, Univ. of Texas M.D. Anderson Cancer Center,  
Science Park, Smithville, TX 78957.

SO Proc Annu Meet Am Assoc Cancer Res, (1996) 37 A804.  
ISSN: 9017-016X.

DT (MEETING ABSTRACTS)

LA English

FS Institute for Cell and Developmental Biology

EM 199701

ED Entered STN: 19980417

Last Updated on STN: 19980417

AB To determine if and when p53 **mutations** occur in the development of squamous **cell** carcinoma (SCC), we studied alterations of this gene in the hamster cheek-pouch carcinogenesis model by using immunohistochemical staining and polymerase chain reaction-single-strand conformation polymorphism (PCR-SSCP) analysis followed by direct DNA sequencing. Twenty four hamsters were treated with 0.5% 7,12-dimethylbenz[a]**anthracene** in mineral oil three times a week for 16 wk. For this study, p53 protein accumulation was evaluated by immunostaining in various hamster cheek pouch lesions including exophytic and endophytic SCC as well as flat dysplastic hyperplasia and carcinomas in situ. A moderate percentage (33.3%) of exophytic lesions was negative for p53 staining, whereas most endophytic carcinomas (90%) showed positive p53 reaction. In addition we also found p53 positive staining in a number of flat lesions including areas of focal hyperplasia, dysplastic hyperplasia, and carcinomas in situ. To determine whether the alterations in p53 staining were due to p53 gene **mutation** we used PCR-SSCP and direct sequencing. PCR products corresponding to exons 5a, 6, 7, and 8 from p53-positive tumors showed shifted bands (four tumors in exon 5, two in exon 6, one in exon 7 and two in exon 8). Direct sequencing of some of the shifted bands revealed three **mutations**. Two of the **mutations** were transversions (G to T) in codons 216 and 252, and the third was a G to C transition in codon 282. This study and additional investigations of this suppressor gene in microdissected lesions may better define the mechanisms of carcinogenesis in the hamster cheek pouch model and may provide a new marker of progression for chemoprevention studies.

L15 ANSWER 197 OF 398 CANCERLIT  
 AN 97609652 CANCERLIT  
 DN 97609652  
 TI The specific N-ras **mutation** in rat  
 7,12-dimethylbenz[a]anthracene (DMBA)-induced leukemia (Meeting  
 abstract).  
 AU Osaka M; Koh T; Matsuo S; Sugiyama T  
 CS Dept. Pathology and Tumor Biology, Postgraduate School of Medicine, Kyoto  
 Univ., Kyoto, 606-01, Japan.  
 SO Non-serial, (1995) Leukemia and Lymphoma, Pathogenesis and Treatment,  
 Molecular Aspects. 18th Symposium of the International Association for  
 Competitive Research on Leukemia and Related Diseases, p. 62. Kyoto,  
 Japan, October 29-November 3, 1995 .  
 DT (MEETING ABSTRACTS)  
 LA English  
 FS Institute for Cell and Developmental Biology  
 EM 199705  
 ED Entered STN: 19980417  
 Last Updated on STN: 19980417  
 AB Intravenous injections of 7,12-dimethylbenz[a]anthracene (DMBA)  
 induce erythroblastic leukemia (erythroleukemia) with 2 trisomy and Long 2  
 in Long-Evans rats. Recently, a consistent type of **mutation**, A  
 to T transversion in codon 61 of N-ras gene, was found in all of 6  
 cultured leukemia **cell** lines and 9 primary leukemias induced by  
 DMBA by polymerase chain reaction (PCR) and direct sequencing (Osaka et  
 al, Cancer Lett; 91:25-31 1995.). On the contrary, no **mutation**  
 was observed in Ha- and Ki-ras genes in all leukemias. The consistent  
 occurrence of above N-ras **mutation** as well as in leukemias  
 indicates that N-ras gene plays an important role in DMBA-leukemogenesis.  
**Mutations** in ras genes are considered to take place during the  
 initiation stage of carcinogenesis because they often appear in the  
 premalignant stage of tumors. In order to detect N-ras **mutation**  
 in early stage of preleukemia, we designed the **mutant**  
 -allele-specific amplification (MASA) method to detect the  
**mutation** in bone marrow (BM) **cells** of DMBA-treated rats.  
 The MASA method was sensitive enough to detect one **mutant** among  
 10(6) normal **cells**. Using this method, the N-ras  
**mutation** was found in BM **cells** 2 days after single DMBA  
 injection and thereafter throughout the preleukemic stage. These results  
 suggest the importance of the N-ras **mutation** as an earliest  
 event in DMBA-leukemogenesis.

L15 ANSWER 196 OF 398           CANCERLIT  
AN 97619054           CANCERLIT  
DN 97619054  
TI Site-specific **mutagenesis** by bulky exocyclic amino-substituted  
guanine and adenine derivatives in E coli and human **cells**.  
(Meeting abstract).  
AU Moon K-Y; Pauly G T; Moschel R C  
CS ABL-BRP, NCI-FCRDC Frederick, MD 21702.  
SO Proc Annu Meet Am Assoc Cancer Res, (1997) 38 A261.  
ISSN: 0197-016X.  
DT (MEETING ABSTRACTS)  
LA English  
FS Institute for Cell and Developmental Biology  
EM 199709  
ED Entered STN: 19980417  
Last Updated on STN: 19980417  
AB The **mutagenicity** of the two major DNA adducts produced by  
7-bromomethylbenz[a]-**anthracene** i.e., N2-(benz[a]anthracen-7-  
ylmethyl)-2'-deoxyguanosine (b[a]a2G) and N6-(benz[a]anthracen-7-ylmethyl)-  
2'-deoxyadenosine (b[a]a6A) as well as the simpler benzylated analogs  
N2-benzyl-2'-deoxyguanosine(bn2G) and N6-benzyl-2'-deoxyadenosine (bn6A)  
was examined in both E. coli and Ad293 human **cells**. In our E.  
coli site-specific **mutagenesis** system none of these aralkylated  
adducts exhibited any significant **mutagenicity** with or without  
SOS induction. In our human **cell** site-specific  
**mutagenesis** system bn2G and bn6A exhibited weak  
**mutagenicity** although b[a]a2G and b[a]a6A were significantly  
**mutagenic**. At the site of the adduct b[a]a2G produced G to T  
transversion **mutations** while b[a]a6A produced A to G transition  
**mutations**. These results indicate that the more bulky b[a]a2G and  
b[a]a6A exhibit significantly greater **mutagenicity** in human  
**cells** than in E. coli and further emphasize the importance of  
studying site-specific **mutagenesis** by carcinogen-modified DNA  
bases in human **cells**.

L15 ANSWER 192 OF 398 MEDLINE

AN 76051128 MEDLINE

DN 76051128 PubMed ID: 1186764

TI Mammalian **cell** transformation and **cell**-mediated **mutagenesis** by carcinogenic polycyclic hydrocarbons.

AU Huberman E

SO MUTATION RESEARCH, (1975 Aug) 29 (2) 285-91.

Journal code: 0400763. ISSN: 0027-5107.

CY Netherlands

DT Journal; Article; (JOURNAL ARTICLE)

LA English

FS Priority Journals

EM 197601

ED Entered STN: 19900313

Last Updated on STN: 19900313

Entered Medline: 19760123

AB The introduction of a polycyclic hydrocarbon such as benzo(alpha)pyrene (BP) into normal golden hamster embryo **cell** cultures results, in addition to cytotoxicity, in malignant **cell** transformation. Studies on the effect of different doses of BP on the normal **cells** showed that the frequency of transformed colonies was directly related to the dose of the carcinogen. Analysis of this dose-response curve suggests a one-event ("one-hit") response for transformation by this carcinogen. The one-event response for transformation by carcinogenic polycyclic hydrocarbons and the fact that these carcinogens bind to DNA in susceptible **cells** suggests that transformation can involve a single alteration in the genetic constitution of the treated **cells**. Carcinogens may, therefore, produce somatic **mutations**, some of which may involve the genes that control malignancy. Recently, considerable progress has been made in developing models for the study of chemical **mutagenesis** in mammalian **cells**. Using resistance to 8-azaguanine as a marker, positive correlations between **mutagenicity** and transformation were obtained with chemically reactive carcinogens such as N-acetoxy-N-2-fluorenyl-acetamide, N-methyl-N'-nitro-N-nitrosoguanidine and K-region epoxides of polycyclic hydrocarbons. However, no such correlations were obtained with the carcinogenic polycyclic hydrocarbons themselves, since the **cell** lines used in chemical **mutagenesis** do not metabolize these carcinogens. In order to obtain better correlations, we have developed a **cell**-mediated **mutagenic** assay with carcinogenic hydrocarbons in which Chinese hamster **cells**, which are susceptible for **mutagenesis**, were co-cultivated with lethally irradiated rodent **cells** that can metabolize these compounds. Using this **cell** mediated assay, we obtained **mutagenesis** with the carcinogenic hydrocarbons 7,12-dimethylbenz(alpha) **anthracene** (DMBA), BP, 3-methylcholanthrene and 7-methylbenz(alpha) **anthracene**; the most potent carcinogen, DMBA, gave the highest frequency of **mutations**. The polycyclic hydrocarbons, pyrene and benz(alpha) **anthracene**, which are not carcinogenic were also not **mutagenic**. We have therefore demonstrated a relationship between the carcinogenicity of polycyclic hydrocarbons and their **mutagenicity** in mammalian **cells**, without having to isolate their reative metabolic intermediates. It should be possible to use in this system human **cells** from different organs and individuals to screen for environmental chemicals hazardous to humans which have to be metabolically activated.

L15 ANSWER 185 OF 398 MEDLINE  
 AN 77206380 MEDLINE  
 DN 77206380 PubMed ID: 873646  
 TI The metabolic activation of 7-methylbenz(a)**anthracene**: the induction of malignant transformation and **mutation** in mammalian **cells** by non-K-region dihydrodiols.  
 AU Marquardt H; Baker S; Tierney B; Grover P L; Sims P  
 SO INTERNATIONAL JOURNAL OF CANCER, (1977 Jun 15) 19 (6) 828-33.  
 Journal code: 0042124. ISSN: 0020-7136.  
 CY Denmark  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 197708  
 ED Entered STN: 19900314  
 Last Updated on STN: 19900314  
 Entered Medline: 19770812  
 AB Four different dihydrodiols derived from 7-methylbenz(a)**anthracene** have been tested, together with the parent hydrocarbon, for their ability to induce the in vitro malignant transformation of **mouse** M2 fibroblasts and **mutations** in V79 Chinese hamster **cells**. In the transformation tests with the non-K-region dihydrodiols, the 3,4-diol was the most active dihydrodiol tested and the 8,9-diol was also more active than 7-methylbenz(a)**anthracene** itself; the 1,2-diol showed only slight activity. The K-region dihydrodiol, the 5,6-diol, which cannot be directly metabolized to a vicinal diol-epoxide, was inactive. These differences in biological activity were similar to those apparent in the results from the **mutagenicity** tests. The data support the general hypothesis that non-I-region dihydrodiols, which can be metabolized to vicinal diol-epoxides, are important in the metabolic activation of the carcinogenic polycyclic hydrocarbons and, when taken together with other results, indicate that 3,4-dihydro-3,4-dihydroxy-7-methylbenz(a)**anthracene** is most probably involved in the metabolic activation of 7-methylbenz(a)**anthracene** presumably following conversion into the related diol-epoxide, 3,4-dihydro-3,4-dihydroxy-7-methylbenz(a)**anthracene** 1,2,-oxide.

L15 ANSWER 186 OF 398 MEDLINE

L15 ANSWER 178 OF 398 MEDLINE  
AN 80176968 MEDLINE  
DN 80176968 PubMed ID: 6768455  
TI Mammary gland **cell**-mediated **mutagenesis** of mammalian  
**cells** by organ-specific carcinogens.  
AU Gould M N  
SO CANCER RESEARCH, (1980 Jun) 40 (6) 1836-41.  
Journal code: 2984705R. ISSN: 0008-5472.  
CY United States  
DT Journal; Article; (JOURNAL ARTICLE)  
LA English  
FS Priority Journals  
EM 198007  
ED Entered STN: 19900315  
Last Updated on STN: 19900315  
Entered Medline: 19800726  
AB **Rat** mammary gland **cells** have been used to activate  
chemical procarcinogens to **mutagenic** compounds in a culture  
system. **Mutagenesis** was tested in Chinese hamster V-79  
**cells** that were cocultured with the mammary **cells**. The  
locus **mutations** tested for were resistance to ouabain and  
resistance to 6-thioguanine. Mammary **cells** were separated into  
several fractions. Fractions enriched in either epithelial or stromal  
**cells** could both mediate **mutagenesis**. The  
**mutation** frequency related to the density of the mammary  
**cells**. The mammary carcinogen 7, 12-dimethylbenz(a)  
**anthracene** exhibited a dose-dependent enhancement of  
**mutation** frequency and cytotoxicity when added to the cocultures,  
whereas the hepatocarcinogen aflatoxin B1 did not. This system may be  
useful in examining some of the mechanisms of organ-specific  
carcinogenesis and also may act as a screening system for carcinogenic  
environmental contaminants.

L15 ANSWER 179 OF 398 MEDLINE

L15 ANSWER 173 OF 398 MEDLINE

AN 82048943 MEDLINE

DN 82048943 PubMed ID: 6271413

TI Lung and liver **cell**-mediated **mutagenesis** systems:  
specificities in the activation of chemical carcinogens.

AU Langenbach R; Nesnow S; Tompa A; Gingell R; Kuszynski C

NC 5-R01-CA20022 (NCI)

SO CARCINOGENESIS, (1981) 2 (9) 851-8.

Journal code: 8008055. ISSN: 0143-3334.

CY United States

DT Journal; Article; (JOURNAL ARTICLE)

LA English

FS Priority Journals

EM 198201

ED Entered STN: 19900316

Last Updated on STN: 19970203

Entered Medline: 19820109

AB A liver and lung **cell**-mediated-V79 **cell**

**mutagenesis** system using intact **cells** as metabolic activation systems was employed to study the relative ability of **cells** from these organs to activate chemical carcinogens. Primary cultures of liver and lung **cells** from male Sprague Dawley rats were used to metabolically activate the chemicals and the **mutation** of Chinese hamster V79 **cells** to ouabain resistance used to detect **mutagenic** intermediates. 7,12-Dimethylbenz[a]anthracene and 3-methylcholanthrene, were more active in the lung system than in the liver **cell** system. Benzo[a]pyrene (B[a]P) was inactive in the liver **cell**-mediated system but **mutagenic** to V79 **cells** in the lung **cell**-mediated system. Dimethylnitrosamine (DMN) was inactive in the presence of liver **cells**. Aflatoxin B1 was **mutagenic** in the liver **cell**-mediated system, but only weakly **mutagenic** in the lung **cell**-mediated system. Because the **mutagenicities** of DMN and B[a]P were organ-specific, the metabolism of these carcinogens in the two primary **cell** systems was investigated. DMN was metabolized by liver but not by lung **cells**, possibly accounting for its lack of **mutagenicity** in the lung **cell** system. B[a]P was extensively metabolized by both **cell** types, but quantitative differences were observed when the metabolic products were analyzed by high pressure liquid chromatography. Comparing total organic and water soluble metabolites, lung **cells** produced similar amounts of 7,8- and 9,10-diols but little 4,5-diol, while liver **cells** produced equivalent total amounts of the three diols. Lung **cells** produced twice the amount of B[a]P glucuronide conjugates as liver **cells**, while liver **cells** produced twice the amount of B[a]P sulfate conjugates as lung. The data suggest that intact **cells** from various organs can be used as metabolic activating systems in vitro assays and that studies into organ specificity can be investigated by this approach.



L15 ANSWER 161 OF 398 MEDLINE  
 AN 83177994 MEDLINE  
 DN 83177994 PubMed ID: 6838477  
 TI The use of DNA-repair-deficient **mutants** of Chinese hamster ovary **cells** in studying **mutagenesis** mechanisms and testing for environmental **mutagens**.  
 AU Thompson L H  
 SO BASIC LIFE SCIENCES, (1983) 23 217-46.  
 Journal code: 0360077. ISSN: 0090-5542.  
 CY United States  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 198305  
 ED Entered STN: 19900318  
 Last Updated on STN: 19900318  
 Entered Medline: 19830505  
 AB Our laboratory has taken a somatic-**cell**-genetics approach to the study of **mutagenesis** by utilizing **mutant** strains of Chinese hamster ovary (CHO) **cells** that are deficient in DNA repair processes. From more than 150 UV-sensitive strains tested, five complementation classes were identified, and representative **mutants** were found to be defective at, or before, the incision step of excision repair. A representative **mutant**, strain UV-5, was compared with the parental strain in terms of cytotoxicity and dose-response curves for **mutation** induction after treatment with UV and several chemicals that are known to produce large adducts in DNA. Excision repair in normal CHO **cells** protects against both cytotoxicity and **mutagenesis**, but the degree of protection depends on both the agent and the genetic marker used for detecting **mutations**. Upon treatment with low doses (100% **cell** survival) of the polyaromatic hydrocarbon 7-bromomethylbenz(a) **anthracene**, repair-deficient UV-5 **cells** had linear responses for **mutation** induction to thioguanine resistance or azaadenine resistance, whereas the normal repair-proficient **cells** showed curvilinear responses in which the slope increased with dose. This behavior suggests that in the normal **cells** the repair system acting on potentially **mutagenic** lesions becomes saturated at doses that produce cytotoxicity. In no instance was a lower **mutation** frequency induced in UV-5 **cells** than the parental **cells**, at a given dose of **mutagen**, suggesting that the excision repair system is error-free in normal CHO **cells**

L15 ANSWER 152 OF 398 MEDLINE  
 AN 84093273 MEDLINE  
 DN 84093273 PubMed ID: 6656796  
 TI Hypersensitivity to **cell** killing and **mutation**  
 induction by chemical carcinogens in an excision repair-deficient  
**mutant** of CHO **cells**.  
 AU Thompson L H; Salazar E P; Brookman K W; Hoy C A  
 SO MUTATION RESEARCH, (1983 Dec) 112 (6) 329-44.  
 Journal code: 0400763. ISSN: 0027-5107.  
 CY Netherlands  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 198402  
 ED Entered STN: 19900319  
 Last Updated on STN: 19900319  
 Entered Medline: 19840214  
 AB A strain of Chinese hamster ovary **cells** that is deficient in  
 nucleotide excision repair, strain UV5, was compared with the normal  
 parental CHO **cells** in terms of cytotoxicity and  
**mutagenesis** after exposure to several chemical carcinogens that  
 are known to produce bulky, covalent adducts in DNA. Induced  
**mutations** were measured at the hprt locus using thioguanine  
 resistance and at the aprt locus using azaadenine resistance. The  
 compounds tested that required metabolic activation (using **rat**  
 or hamster microsomal fractions) were 7,12-dimethylbenz(a)  
**anthracene**, 3-methylcholanthrene, benzo(a)pyrene, aflatoxin B1,  
 2-acetylaminofluorene, and 2-naphthylamine. The direct-acting compounds  
 (+/-)-r-7,t-8-dihydroxy-t-9,10-epoxy-7,8,9,10-tetrahydrobenzo(a)pyrene,  
 N-acetoxy-2-acetylaminofluorene, and N-OH-2-naphthylamine were also  
 studied. For all compounds except 2-naphthylamine and its active  
 metabolite, the repair-deficient **cells** were significantly more  
 sensitive to killing than the normal CHO **cells**. **Mutation**  
 induction at both loci was also more efficient in UV5 **cells** in  
 each instance where enhanced cytotoxicity was observed. By using  
 tritium-labeled N-acetoxy-2-acetylaminofluorene, normal and **mutant**  
**cells** were shown to bind **mutagen** to their nuclear DNA  
 with similar efficiency, and a greater amount of adduct removal occurred  
 in the normal **cells**. From this study it is concluded that the  
 use of excision repair-deficient CHO **cells** provides enhanced  
 sensitivity for detecting **mutagenesis** and that a positive  
 differential cytotoxicity response gives an indication of repairable,  
 potentially lethal genetic damage.

L15 ANSWER 123 OF 398 MEDLINE  
 AN 88079999 MEDLINE  
 DN 88079999 PubMed ID: 3121168  
 TI Development of **murine** epidermal **cell** lines which  
 contain an activated rasHa oncogene and form papillomas in skin grafts on  
 athymic nude **mouse** hosts.  
 AU Strickland J E; Greenhalgh D A; Koceva-Chyla A; Hennings H; Restrepo C;  
 Balaschak M; Yuspa S H  
 CS Laboratory of Cellular Carcinogenesis and Tumor Promotion, National Cancer  
 Institute, Bethesda, Maryland 20892.  
 SO CANCER RESEARCH, (1988 Jan 1) 48 (1) 165-9.  
 Journal code: 2984705R. ISSN: 0008-5472.  
 CY United States  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 198802  
 ED Entered STN: 19900305  
 Last Updated on STN: 19900305  
 Entered Medline: 19880208  
 AB We have developed four **murine** epidermal **cell** lines  
 which form squamous papillomas when grafted to athymic nude mice in a  
 reconstituted skin. Two of the lines, SP-1 and BP-4, were derived from  
 pools of papillomas produced on SENCAR and BALB/c **mouse** skin,  
 respectively, by initiation with 7,12-dimethylbenz(a)**anthracene**  
 and promotion with 12-O-tetradecanoylphorbol-13-acetate. Line 308 was  
 derived from BALB/c **mouse** skin initiated in **vivo** with  
 7,12-dimethylbenz(a)**anthracene**, culture of the epidermal  
**cells**, and selection of **cells** resistant to Ca2+-induced  
 terminal differentiation. Line LC14 was derived from untreated, cultured  
 newborn BALB/c **mouse** primary epidermal **cells** which  
 spontaneously developed resistance to Ca2+-induced terminal  
 differentiation. Each line has an activated rasHa gene with a  
**mutation** within codon 61. **Cells** from all four lines, in  
 contrast to normal primary epidermal **cells**, survive in medium  
 with Ca2+ levels greater than 0.1 mM. Clonal growth studies in culture  
 showed a unique growth pattern for each of the four lines in medium with  
 1.4 mM and 0.05 mM Ca2+, with or without 12-O-tetradecanoylphorbol-13-  
 acetate. Early passage **cells** of these lines should provide a  
 valuable resource for detecting genes or genetic alterations which  
 complement an activated ras gene to cause malignant conversion and for  
 studying the biology of tumor promotion.

tumor and suggest that it could be involved in tumor regression.

L15 ANSWER 104 OF 398 MEDLINE  
AN 91058635 MEDLINE  
DN 91058635 PubMed ID: 1978778  
TI Ha-ras oncogene **mutations** in **cell** lines derived from  
**rat** tracheal implants exposed in **vivo** to  
7,12-dimethylbenz[a]**anthracene**.  
AU Cosma G N; Wirgin I I; Marchok A C; Garte S J  
CS Institute of Environmental Medicine, New York University Medical Center,  
New York 10016.  
NC CA13343 (NCI)  
CA36342 (NCI)  
CA42798 (NCI)  
+  
SO MOLECULAR CARCINOGENESIS, (1990) 3 (5) 258-63.  
Journal code: 8811105. ISSN: 0899-1987.  
CY United States  
DT Journal; Article; (JOURNAL ARTICLE)  
LA English  
FS Priority Journals  
EM 199101  
ED Entered STN: 19910222  
Last Updated on STN: 20000303  
Entered Medline: 19910110  
AB The frequency of Ha-ras **mutations** was determined as a function  
of neoplastic progression in **cell** lines derived from **rat**  
tracheal implants exposed in **vivo** to 7,12-dimethylbenz[a]  
**anthracene**. Restriction fragment-length polymorphism (RFLP)  
analysis revealed an A---T transversion in the second base of codon 61 in  
2 of 11 **cell** lines. One of the positive **cell** lines was  
tumorigenic, but the other was neither tumorigenic nor anchorage  
independent, thus indicating a lack of correlation between neoplastic  
stage and ras **mutation**. Densitometry analysis of the RFLP bands  
indicated that approximately 50% of the **cells** within these two  
heterogeneous populations contained the **mutation**. Direct  
sequence analysis of polymerase chain reaction-amplified DNA confirmed  
these results and did not reveal any other **mutations** in this  
region of the Ha-ras gene.

L19 ANSWER 1 OF 6 EMBASE COPYRIGHT 2002 ELSEVIER SCI. B.V.

AN 93208547 EMBASE

DN 1993208547

TI 1,2-Dimethyl-9, 10 benzantracene induced  
malignant fibrous histiocytoma in rats and the effect of adrenalectomy on  
tumor growth.

AU Kotiloglu E.; Oner U.

CS Pediatric Pathology Unit, Hacettepe University Medical School, Ankara,  
Turkey

SO Doga - Turkish Journal of Medical Sciences, (1993) 18/2 (115-126).  
ISSN: 1010-7584 CODEN: DTJSEX

CY Turkey

DT Journal; Article

FS 005 General Pathology and Pathological Anatomy

016 Cancer

052 Toxicology

LA English

SL English

L19 ANSWER 2 OF 6 CAPLUS COPYRIGHT 2002 ACS

AN 1993:249623 CAPLUS

DN 118:249623

TI Comparison of target organs of carcinogenicity for mutagenic and  
non-mutagenic chemicals

AU Gold, Lois Swirsky; Slone, Thomas H.; Stern, Bonnie R.; Bernstein, Leslie

CS Life Sci. Div., Lawrence Berkeley Lab., Berkeley, CA, 94720, USA

SO Mutat. Res. (1993), 286(1), 75-100

CODEN: MUREAV; ISSN: 0027-5107

DT Journal

LA English

AB A comparison of target organs for mutagens and non-mutagens is presented  
for 351 rodent carcinogens in the Carcinogenic Potency Database (CPDB)  
with mutagenicity evaluations in Salmonella. Results are consistent with  
the hypotheses that in high-dose rodent tests mitogenesis is important in  
the carcinogenic response for mutagens and non-mutagens alike, and that  
mutagens have a multiplicative interaction for carcinogenicity because  
they can both damage DNA directly and cause **cell** division at  
high doses. Among carcinogens that induce tumors at multiple sites in  
both rats and mice, 81% are mutagens; in comparison, among carcinogens  
that are pos. at only a single target site in one species and are neg. in  
the other, 42% are mutagens. Both mutagens and non-mutagens induce tumors  
in a wide variety of sites, and most organs are target sites for both.  
Moreover, the same sites tend to be the most common sites for both: 79% or  
more of both mutagenic and non-mutagenic carcinogens are pos. in each  
species in at least one of the 8 most frequent target sites: liver, lung,  
mammary gland, stomach, vascular system, kidney, hematopoietic system and  
urinary bladder. Species differences are discussed as well as results for  
particular target organs: liver, Zymbal's gland and kidney. A compendium  
of bioassay results is presented.

L21 ANSWER 46 OF 56 BIOSIS COPYRIGHT 2002 BIOLOGICAL ABSTRACTS INC.  
 AN 2001:276690 BIOSIS  
 DN PREV200100276690  
 TI In vitro and in **vivo** chemopreventive properties of a soybean peptide (lunasin).  
 AU Lam, Yi (1); Chen, Na (1); de Lumen, Benito (1)  
 CS (1) University of California, Berkeley, CA USA  
 SO FASEB Journal, (March 7, 2001) Vol. 15, No. 4, pp. A281. print.  
 Meeting Info.: Annual Meeting of the Federation of American Societies for Experimental Biology on Experimental Biology 2001 Orlando, Florida, USA March 31-April 04, 2001  
 ISSN: 0892-6638.  
 DT Conference  
 LA English  
 SL English  
 AB Lunasin is a non-abundant 43-amino acid peptide found in soybean and contains an -RGD-**cell** adhesion motif followed by 8 aspartic acid (D) residues at the carboxyl end. Previous studies have demonstrated the ability of lunasin to bind to chromatin, leading to disruption of kinetochore formation and inhibition of mitosis when the lunasin **gene** is transfected into mammalian **cells**. Experiments reported here indicate that lunasin peptide can inhibit carcinogenesis both in **vivo** and > in vitro. We studied the effect of topically applied lunasin in a two-stage protocol of skin carcinogenesis in female SENCAR mice. At 13 weeks of age mice were initiated with 5 mug 7,12-dimethylbenz(a)**anthracene** (DMBA), and promoted with 2 mug 12-O-tetradecanoylphorbol-13-acetate (TPA) twice per week for 19 weeks. Starting from the week prior to DMBA administration till the end of the study, mice were subjected to 2.5 mug, 25 mug or 250 mug of lunasin peptide applied topically in 100% ethanol weekly. A significant reduction in tumor incidence and tumor latency was observed in the high-dosage (250 mug/week) lunasin group but not in the low (2.5 mug/week) and medium (2.5 mug/week) dosage groups. These results demonstrate that high dosage of lunasin is effective in inhibiting skin carcinogenesis and this effect could have been exerted during the initiation or the promotion stages. Lunasin peptide has also been shown to suppress ras induced transformation of **mouse** fibroblast NIH 3T3 **cells** in vitro by 40% at a concentration as low as 100nM. Lunasin had an irreversible effect on ras-transformation process since treatment with lunasin for 3 days was effective in suppressing transformation. Lunasin can suppress foci formation even when added 7 days after ras transfection of 3T3 **cells**. Using various deletion **mutant** forms of lunasin in the foci **assay**, we found that the polyaspartic end of lunasin is necessary for the transformation suppression effect. Furthermore lunasin reduced anchorage independent growth of stably ras-transfected **mouse** fibroblast **cells** by 40% in a colony formation **assay**. In addition, Western analyses show that lunasin increases p21 expression in ras-transfected **cells**.

-12345X@PJL

@PJL JOB NAME = "MSJOB 57"

@PJL USTATUS JOB = ON

@PJL USTATUS PAGE = OFF

@PJL USTATUS DEVICE = ON

@PJL USTATUS TIMED = 30

L21 ANSWER 28 OF 56 EMBASE COPYRIGHT 2002 ELSEVIER SCI. B.V.  
 AN 2001158665 EMBASE  
 TI Promoting effect of a high-fat/high-protein diet in DMBA-induced ductal pancreatic cancer in rats.  
 AU Z'graggen K.; Warshaw A.L.; Werner J.; Graeme-Cook F.; Jimenez R.E.; Fernandez-del Castillo C.; Urist M.M.; Townsend C.M. Jr.; Warshaw A.L.  
 CS Dr. C. Fernandez-del Castillo, Department of Surgery, Massachusetts General Hospital, WACC 336, Boston, MA 02114, United States  
 SO Annals of Surgery, (2001) 233/5 (688-695).  
 Refs: 21  
 ISSN: 0003-4932 CODEN: ANSUA5  
 CY United States  
 DT Journal; Conference Article  
 FS 009 Surgery  
 016 Cancer  
 029 Clinical Biochemistry  
 LA English  
 SL English  
 AB Objective: To investigate whether a high-fat/high-protein diet (HFPD) acts as a promoter of the natural course of cancer growth in the 7,12-dimethylbenzanthracene (DMBA)-induced ductal pancreatic cancer model in rats. Summary Background Data: DMBA implantation to the **rat** pancreas induces ductal adenocarcinoma. Information regarding the effects of diet and the presence of K-ras **mutation** in this model is not available. Methods: Rats were randomly assigned to regular **rat** chow or a diet with a 30% content in fat and protein (HFPD). The presentation of cancer, the histologic spectrum of neoplasia at 1 and 9 months, and the prevalence of cancer in relation to diet were assessed. Histologic specimens comprising normal ducts, hyperplasia, dysplasia/carcinoma in situ, or carcinoma were designated by a pathologist and microdissected. Genomic DNA was extracted, and K-ras and H-ras **gene mutations** were determined by a **mutant** -enriched polymerase chain reaction **assay** and direct sequencing. Results: Rats fed HFPD increased their weight significantly compared with controls. DMBA induced characteristic stages of neoplasia at the implant site but not elsewhere. Macroscopic cancers of the pancreatic head presented regularly with common bile duct and gastric outlet obstruction. The prevalence of K-ras **mutations** was proportional to the degree of epithelial abnormality. K-ras **mutations** were significantly more frequent in cancer than in normal and hyperplastic ducts. H-ras **mutations** were not found. At 1 month in the HFPD-fed rats, the prevalence of cancer (16%) and dysplasia (16%) was not significantly different from the prevalence of cancer (29%) and dysplasia (8%) in the chow-fed rats. At 9 months the prevalence of cancer in the HFPD-fed rats increased to 29%, whereas that in the chow-fed rats decreased to 17%. The combined prevalence of cancer and dysplasia at 9 months in the HFPD-fed rats (34%) significantly exceeded that in the chow-fed rats. Conclusions: DMBA induces characteristic stages of neoplasia in the evolution of ductal pancreatic cancer in rats. K-ras **mutations** occur progressively in the ladder of oncogenesis, as in human pancreatic neoplasms. The addition of a diet with a high fat and protein content acts as a promoter of carcinogenesis, possibly by interfering with repair mechanisms and natural regression of early lesions.



L21 ANSWER 26 OF 56      CANCERLIT

AN 88643618      CANCERLIT

DN 88643618

TI ACTIVATION OF RAS ONCOGENES BY CHEMICAL CARCINOGENS.

AU Barbacid M; Sukumar S; Zarbl H

CS Developmental Oncology Section, LBI-Basic Res. Program, NCI-Frederick  
Cancer Res. Facility, P.O. Box B, Building 539, Frederick, MD.

SO Gene Amplification Anal, (1986) 4 21-38.

ISSN: 0275-2778.

DT Journal; Article; (JOURNAL ARTICLE)

LA English

FS Institute for Cell and Developmental Biology

EM 198805

ED Entered STN: 19941107

Last Updated on STN: 19950508

AB Data from the authors' and others' laboratories pertaining to the mechanism of activation and involvement of ras oncogenes in human and carcinogen-induced animal tumors are reviewed. High proportions (79% and 72%, respectively) of N-nitroso-N-methylurea (NMU)-induced mammary carcinomas contained H-ras-1 oncogenes in experiments with Buf/N or Sprague-Dawley rats. A polymorphic Mnl I DNA fragment was generated and used in a **gene transfer assay** for the presence of **mutated H-ras-1 gene**. Altogether, 61/71 NMU-induced tumors scored positive in the Mnl I **assay**, including 32/61 for Buf/N and 23/61 for Sprague-Dawley. Of these 55 mammary carcinomas known to contain H-ras-1 oncogenes by molecular assays, only 48 (30 Buf/N and 18 Sprague-Dawley) scored positive in **gene transfer assays**. These observations suggest a close association between **mutagenesis** in this specific polymorphic Mnl I fragment of the H-ras-1 locus and tumor development. Nonadecamers were generated that were capable of identifying substitutions in position 35 (specific point **mutations**) in genomic DNA. This residue is the only nucleotide of the Mnl I cleavage site that can alter the coding properties of the critical twelfth codon of the H-ras-1 **gene**. With this system, the presence of activating deoxyguanosine to deoxyadenosine or deoxythymidine **mutations** of H-ras-1 oncogenes was demonstrated unequivocally in DNA isolated from tumor tissues. Using a nonadecamer probe complementary to the 61st codon (CAA) and flanking sequences of the normal **rat** H-ras-1 locus, it was shown that each of the five H-ras-1 oncogenes present in 7,12-dimethylbenz(a)**anthracene**-induced mammary carcinomas harbored **mutations** in the region of the 61st codon. It was further demonstrated that the G to A **mutations** present in each of the NMU-induced H-ras-1 oncogenes do not result from either positive growth selection or specific repair systems. Instead, the malignant activation of the H-ras-1 locus in NMU-induced mammary carcinomas results from the direct **mutagenic** effect of NMU on this locus. It is very likely that some environmental carcinogens might initiate human tumors by activating ras oncogenes. (50 Refs)

L21 ANSWER 24 OF 56      CANCERLIT

AN 94697099      CANCERLIT

DN 94697099

TI N-methyl-N-nitrosourea induced **rat** mammary tumors arise from  
**cells** harboring spontaneous oncogenic Ha-ras-1 **gene**  
**mutations.**

AU Cha R S

CS Massachusetts Institute of Technology.

SO Diss Abstr Int [B], (1993) 54 (3) 1363.

ISSN: 0419-4217.

DT (THESIS)

LA English

FS Institute for Cell and Developmental Biology

EM 199405

ED Entered STN: 19941107

Last Updated on STN: 19970509

AB The overall goal of this project was to directly measure the number of  
**cells** harboring the specific oncogenic **mutations** in  
target tissues of animals before and after carcinogen exposure. The  
NMU-induced **rat** mammary tumor model was chosen. The direct  
measurements of the specific ras **mutants** in animal tissues  
required the development of a polymerase chain reaction (PCR)-based  
procedure (mismatch amplification **mutation assay** or  
MAMA) which could detect the G-to-A transitions at the 12th codon of the  
Ha-ras **gene** present at a frequency of 10(-5). This procedure was  
then applied in measuring the number of specific ras **mutants** in  
the mammary epithelium before and after NMU exposure. Based on these  
results, it was concluded that the NMU-induced mammary tumors carrying the  
specific Ha-ras **gene mutations** arose from pre-existing  
ras **mutants**. Presumably, an independent effect(s) of NMU in one  
of these **mutant cells** was responsible for tumor  
formation. In an attempt to investigate molecular mechanisms underlying  
tissue specificity in NMU-induced tumorigenesis, MAMA was also carried out  
on the nontarget liver tissue before and after NMU exposure. These results  
were then compared to the results obtained from mammary tissue. These  
observations suggest that the lower **mutation** rate for the  
specific G-to-A transitions of the Ha-ras **gene** and the fact that  
the ras **mutants** do not acquire a significant growth advantage in  
the liver tissue may contribute to the apparent resistance of liver tissue  
to NMU-induced tumorigenesis. Molecular analysis of NMU- and  
dimethylbenz(a)**anthracene** (DMBA)-induced, and spontaneously  
arising mammary lesions was carried out utilizing denaturant gradient gel  
electrophoresis. None of the 31 spontaneously arising mammary lesions  
carried activated Ha-ras genes. These results suggest that the  
pre-existing ras **mutants** are specifically promoted during  
NMU-induced tumorigenesis, but not during DMBA-induced, or spontaneous  
tumorigenesis. The conspicuous absence of this G-to-A **mutation**  
among DMBA-induced and spontaneously arising mammary lesions suggest that  
independent molecular mechanisms are responsible for development of  
mammary lesions in these systems. (Copies available exclusively from MIT  
Libraries, Rm. 14-0551, Cambridge, MA 02139-4307. Ph. 617-253-5668; Fax  
617-253-1690. Abstract shortened by UMI. Not available from University  
Microfilms Int'l.)

L21 ANSWER 23 OF 56      CANCERLIT  
 AN 96604700      CANCERLIT  
 DN 96604700  
 TI Carcinogen induced mechanisms in mammary tumorigenesis (Meeting abstract).  
 AU Jin Z; Zarbl H  
 CS Fred Hutchinson Cancer Research Center, Seattle, WA 98104-2092.  
 SO Proc Annu Meet Am Assoc Cancer Res, (1995) 36 A658-9.  
 ISSN: 0197-016X.  
 DT (MEETING ABSTRACTS)  
 LA English  
 FS Institute for Cell and Developmental Biology  
 EM 199605  
 ED Entered STN: 19970509  
 Last Updated on STN: 19970509  
 AB We developed a Mismatch Amplification **Mutation Assay** (MAMA) capable of detecting codon twelve GGA to GAA **mutations** in the Ha-ras-1 **gene** when present at a frequency of one in 10(5) alleles. The MAMA was used to measure the frequency of the activating Ha-ras-1 **gene mutation** in mammary epithelial **cells** (RMECs) of 50 day old, virgin, female Fischer 344 rats, before and after exposure to a single carcinogenic dose of N-nitroso-N-methylurea (NMU). The codon twelve G to A transitions arose as background **mutations** in the developing mammary gland and Ha-ras **mutants** were clustered within organ sectors, consistent with their origin during mammary gland development. Exposure to a single carcinogenic dose of NMU failed to induce a significant increase in the number of Ha-ras-1 **mutants**, the fraction of organ sectors containing **mutant cells**, or the fraction of animals harboring mammary epithelial **cells** with Ha-ras-1 **gene mutations**. Thus, the vast majority of NMU-induced carcinomas arise from mammary epithelial **cells** which harbored Ha-ras-1 oncogenes prior to carcinogen exposure. Our results further indicated that NMU allows for the outgrowth of pre-existing Ha-ras-1 **mutants**, and that one or more of these clones eventually gives rise to mammary carcinomas. Although the target for NMU-induced **mutagenesis** could be composed of a large set of cooperating oncogenes or tumor suppressor genes, it was equally plausible that the carcinogenic effect of NMU is mediated via nonmutagenic mechanisms. We used Southern blot analyses of genomic DNAs digested with either the HpaII or MspI restriction enzymes (methylation sensitive and insensitive isoschizomers, respectively) to detect changes in Ha-ras-1 **gene** DNA methylation after treatment of rats with NMU. While these studies failed to detect any NMU-induced changes in Ha-ras-1 **gene** methylation, they did reveal the presence of a HpaII/MspI restriction site within the Ha-ras-1 promoter region of DNA isolated from RMECs which was refractory to digestion by either enzyme. The same site could however be digested with MspI in DNA isolated from liver. We developed a Polymerase Chain Reaction (PCR)-based **assay** which allows us to determine whether the Ha-ras-1 **gene** promoters in DNA isolated from RMECs or liver **cells** were differentially sensitive to agents which preferentially cleave single-stranded DNA. Genomic DNAs isolated from RMECs or liver were treated with mung bean nuclease or potassium permanganate. The ability of these single-strand specific reagents to cleave the Ha-ras-1 promoter region was then assessed by measuring the efficiency with which PCR primers flanking the HpaII/MspI site in question amplified a 439 bp fragment. The results of these analyses indicated that single-strand specific reagents abrogated amplification from RMECs DNA under treatment conditions that allowed amplification of liver DNA with an efficiency of about 70% per cycle efficiency. These results are consistent with the hypothesis that in RMECs, a region of the Ha-ras promoter is involved in a topological feature (toposwitch) which renders the DNA refractory to

digestion by the HpaII and MspI enzymes. In addition to RMECs, the toposwitch was also detected in a subpopulation of lung and spleen **cells**, but not in liver or kidney **cells**. More importantly however, the loss of the toposwitch appears to play a role in NMU-induced mammary carcinogenesis. The region of the promoter containing the toposwitch became sensitive to MspI digestion in 100% of NMU induced mammary tumors, but remained refractory to digestion in 70% of DMBA-induced mammary tumors. Exposure of pubescent female rats to a carcinogenic dose of NMU initiates the in **vivo** loss of this tissue specific DNA conformation (toposwitching) in greater than 90% of RMECs of exposed animals, with a half life of about seven days. Exposure to a carcinogenic dose of dimethylbenz(a)**anthracene** (DMBA) failed to induce a detectable amount of toposwitching even at 30 days after exposure, suggesting that the latter occurs as a late spontaneous event in 30% of DMBA-indu(ABSTRACT TRUNCATED)

L21 ANSWER 22 OF 56 CANCERLIT  
 AN 96649123 CANCERLIT  
 DN 96649123  
 TI The detection of **gene mutation** in transgenic mice (**Muta Mouse**) following administration of known **mutagens** (Meeting abstract).  
 AU Brooks T M; Dean S W; Kirkland D J  
 CS Hazleton Europe Limited, Otley Road, Harrogate, North Yorkshire HG3 1PY, UK.  
 SO Environ Mol Mutagen, (1995) 25 (Suppl 25) 6.  
 ISSN: 0893-6692.  
 DT (MEETING ABSTRACTS)  
 LA English  
 FS Institute for Cell and Developmental Biology  
 EM 199608  
 ED Entered STN: 19970509  
 Last Updated on STN: 19970509  
 AB We are currently validating the **Muta Mouse** positive selection (lacZ/gale) **assay** to detect **mutation** in a tissue using known **mutagens**/carcinogens. 2-acetylaminofluorene (2-AAF) was administered as a single oral dose at 50 or 100 mg/kg and mice sacrificed 3, 7, 14, 28, 56 or 112 days after treatment. Cyclophosphamide (CPA) was dosed orally at 5 x 40 or 80 mg/kg and sampled at similar intervals. **Mutation** frequencies were determined in DNA from liver following 2-AAF and from bone marrow following CPA treatment. A **mutagenic** response was observed in mice treated at 100 mg/kg 2-AAF, seen from 28 up to 112 days after the single exposure. A small effect was seen in bone marrow following 5 x 80 mg/kg CPA treatment only at 3 days. Animals were treated with a single oral dose of N-methyl-N'-nitro-N-nitrosoguanidine (MNNG) at 50 or 100 mg/kg or 1-chloromethylpyrene (CMP) at 25 or 50 mg/kg, and sampled 3, 7 and 10 days after treatment. Marked dose- and time-dependent increases in **mutation** frequency were seen in DNA from stomachs of MNNG-treated animals but not in CMP-treated mice. Other animals were treated topically by a single application, in acetone, of 250 or 500 ug MNNG, 5 or 10 ug CMP or 40 ug 7,12-dimethylbenz[a]anthracene (DMBA). Both MNNG and DMBA caused marked increases in the **mutation** frequency in DNA from treated skin, whereas only a small effect was seen with CMP. These data further demonstrate the potential of such **assay** systems for the measurement of **gene mutation** induced in **vivo** by direct and indirect-acting **mutagens**.

L21 ANSWER 17 OF 56 MEDLINE  
 AN 91000223 MEDLINE  
 DN 91000223 PubMed ID: 2119594  
 TI Relationship between chemically induced Ha-ras **mutation** and transformation of BALB/c 3T3 **cells**: evidence for chemical-specific activation and **cell** type-specific recruitment of oncogene in transformation.  
 AU Nakazawa H; Aguelon A M; Yamasaki H  
 CS International Agency for Research on Cancer, Lyon, France.  
 NC R01 CA40534 (NCI)  
 SO MOLECULAR CARCINOGENESIS, (1990) 3 (4) 202-9.  
 Journal code: 8811105. ISSN: 0899-1987.  
 CY United States  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 199011  
 ED Entered STN: 19910117  
 Last Updated on STN: 19970203  
 Entered Medline: 19901105  
 AB BALB/c 3T3 **cells** were exposed to 7,12-dimethylbenz[a]anthracene (DMBA) and resultant transformed foci were analyzed for the presence of A182----T **mutation** at codon 61 of Ha-ras (a **mutation** found in many DMBA-induced animal tumors). None of the 30 independently cloned transformed **cell** lines contained such a **mutation**. In order to see whether DMBA is able to induce this **mutation** in BALB/c 3T3 **cells**, we developed a method sensitive enough to detect this specific **mutation** at the frequency of  $10^{-6}$ . Employing this **assay**, we found time- and dose-dependent induction by DMBA of Ha-ras A182----T **mutation** in BALB/c 3T3 **cells**; for example, 2 wk after exposure to 100 micrograms/mL DMBA, 1.4 in  $1 \times 10^4$  **cells** contained this specific **mutation**. On the other hand, other agents that also induce BALB/c 3T3 **cell** transformation, such as 3-methylcholanthrene (MCA), 12-O-tetradecanoylphorbol-13-acetate (TPA), N-methyl-N'-nitro-N-nitrosoguanidine (MNNG), or ultraviolet light, did not induce the **mutation** at detectable frequency (less than  $10^{-6}$ ). These results suggest that DMBA efficiently induces Ha-ras **mutation** in BALB/c 3T3 **cells** but that this **mutation** is not recruited in the process of **cell** transformation. A hypothesis of carcinogen-specific **mutation** of Ha-ras **gene** and its tissue (**cell** type)-specific recruitment in carcinogenesis is proposed.

L21 ANSWER 15 OF 56 MEDLINE  
 AN 92335305 MEDLINE  
 DN 92335305 PubMed ID: 1352887  
 TI Detection of **mutant** Ha-ras genes in chemically initiated **mouse** skin epidermis before the development of benign tumors.  
 CM Erratum in: Proc Natl Acad Sci U S A 1993 Jan 15;90(2):781  
 AU Nelson M A; Futscher B W; Kinsella T; Wymer J; Bowden G T  
 CS Department of Radiation Oncology, College of Medicine, University of Arizona, Tucson 85724.  
 NC CA 40584 (NCI)  
 ES05533-01 (NIEHS)  
 SO PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF AMERICA, (1992 Jul 15) 89 (14) 6398-402.  
 Journal code: 7505876. ISSN: 0027-8424.  
 CY United States  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 199208  
 ED Entered STN: 19920904  
 Last Updated on STN: 19950206  
 Entered Medline: 19920818  
 AB An activated Ha-ras oncogene has been consistently found in chemically initiated benign and malignant **mouse** skin tumors, and an activated ras oncogene has been shown to initiate the process of **mouse** skin carcinogenesis. However, the exact timing of **mutational** activation of the Ha-ras **gene** relative to application of the chemical carcinogen is not known. A sensitive **mutation**-specific PCR technique was used to experimentally address the timing of Ha-ras **gene mutational** activation. This technique can detect **mutant** Ha-ras alleles in the presence of a very large excess of normal ras alleles. Activated Ha-ras genes with 61st codon A----T **mutations** were found in the epidermis of mice 1 week after topical initiation with 7,12-dimethylbenz[a]**anthracene** or urethane by using this **assay**. These results were confirmed by Xba I restriction fragment length polymorphism analysis and direct DNA sequencing. One week after initiation is 1-2 months before the appearance of benign papillomas that harbor activated Ha-ras oncogenes when the initiated mice are promoted with the tumor promoter phorbol 12-myristate 13-acetate. Our data support the hypothesis that initiated epidermal **cells** containing an activated Ha-ras **gene** can remain dormant in the skin until a tumor promoter induces regenerative hyperplasia that allows for outgrowth of these **cells** with an activated ras oncogene to give rise to a benign papilloma.

L21 ANSWER 13 OF 56 MEDLINE  
AN 93196513 MEDLINE  
DN 93196513 PubMed ID: 8450770  
TI Detection of chemical **mutagens** using **Muta**  
**Mouse**: a transgenic **mouse** model.  
AU Hoorn A J; Custer L L; Myhr B C; Brusick D; Gossen J; Vijg J  
CS Hazleton Washington, Inc, Vienna, VA 22182.  
SO MUTAGENESIS, (1993 Jan) 8 (1) 7-10.  
Journal code: 8707812. ISSN: 0267-8357.  
CY ENGLAND: United Kingdom  
DT Journal; Article; (JOURNAL ARTICLE)  
LA English  
FS Priority Journals  
EM 199304  
ED Entered STN: 19930423  
Last Updated on STN: 19930423  
Entered Medline: 19930415  
AB A transgenic **mouse** strain with a high copy number of rescuable lacZ sequences was evaluated for its effectiveness in detecting lacZ-**mutations** in selected tissues. Procarbazine, cyclophosphamide, ethylnitrosourea, 7,12-dimethylbenz[a]**anthracene** (DMBA), acrylamide and chlorambucil were tested following either single or repeated dosing regimens. Bone marrow, liver, skin and testis tissues were selected to assess as target sites for **mutation**. Bone marrow, liver and testis tissues were examined for **mutation** following exposures to ethylnitrosourea and chlorambucil. Increased **mutant** frequencies were found for both chemicals in all three tissues. Bone marrow tissue was examined for **mutation** following procarbazine, cyclophosphamide and acrylamide exposures, and skin was examined for **mutation** following dermal application of DMBA. **Mutation** induction was observed in all cases. The results obtained from this investigation demonstrate the applicability of this transgenic **mouse** as an effective model to detect and analyze **gene mutation** in selected organs including germinal tissues. Studies of organotrophic chemical **mutagens** and carcinogens are possible with this model as are studies of the susceptibility of germinal tissues to **mutagen** exposures.



L21 ANSWER 12 OF 56 MEDLINE  
 AN 95285522 MEDLINE  
 DN 95285522 PubMed ID: 7767973  
 TI Dose-related changes in the profile of ras **mutations** in chemically induced CD-1 **mouse** liver tumors.  
 AU Manam S; Shinder G A; Joslyn D J; Kraynak A R; Hammermeister C L; Leander K R; Ledwith B J; Prahalada S; van Zwieten M J; Nichols W W  
 CS Department of Safety Assessment, Merck Research Laboratories, West Point, PA 19486, USA.  
 SO CARCINOGENESIS, (1995 May) 16 (5) 1113-9.  
 Journal code: 8008055. ISSN: 0143-3334.  
 CY ENGLAND: United Kingdom  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 199507  
 ED Entered STN: 19950713  
 Last Updated on STN: 19950713  
 Entered Medline: 19950705  
 AB We investigated the role of dosing regimen on ras **mutations** in chemically induced CD-1 **mouse** liver tumors. The spectra of ras **gene mutations** in liver tumors that were induced by 15 daily i.p. injections of 7,12-dimethylbenz[a]**anthracene** (DMBA), 4-aminoazobenzene (AAB), N-hydroxy-2-acetylaminofluorene (N-OH-AAF) or N-nitrosodiethylamine (DEN) were compared to those previously obtained for tumors induced by a single but higher dose of each carcinogen. The principal **assay** used was a direct tumor analysis involving sequencing of polymerase chain reaction (PCR)-amplified tumor DNA; additional **mutations** that were present in only a small fraction of tumor **cells** were detected using a transfection **assay** or a PCR-engineered restriction fragment length polymorphism method. Spontaneous liver tumors had a relatively low frequency of ras **mutations**, all found in Ha-ras codon 61, and most of these **mutations** were present in only a small fraction of tumor **cells**. With the exception of multiple-dose DEN, each group of single- and multiple-dose carcinogen-induced tumors exhibited a higher frequency of ras **mutations** compared with spontaneous tumors. For AAB, N-OH-AAF and DEN, the dosing regimen was found to affect significantly the profile of ras **mutations**. For each of these carcinogens, the multiple-dose tumor group (versus single-dose group) had fewer Ki-ras and N-ras **mutations** and more tumors in which the Ha-ras codon 61 (C-->A) **mutation** was present in a large fraction of **cells**. Our results demonstrate that the dosing procedure can materially affect the pattern of ras **gene mutation** in **mouse** liver tumors.

L21 ANSWER 8 OF 56 MEDLINE  
 AN 1999452907 MEDLINE  
 DN 99452907 PubMed ID: 10521810  
 TI A functional and quantitative **mutational** analysis of p53  
**mutations** in yeast indicates strand biases and different roles of  
**mutations** in DMBA- and BBN-induced tumors in rats.  
 CM Erratum in: Int J Cancer 2000 Mar 15;85(6):898  
 AU Yamamoto K; Nakata D; Tada M; Tonoki H; Nishida T; Hirai A; Ba Y; Aoyama  
 T; Hamada J; Furuuchi K; Harada H; Hirai K; Shibahara N; Katsuoka Y;  
 Moriuchi T  
 CS Division of Cell Biology, Hokkaido University School of Medicine, Sapporo,  
 Japan.  
 SO INTERNATIONAL JOURNAL OF CANCER, (1999 Nov 26) 83 (5) 700-5.  
 Journal code: 0042124. ISSN: 0020-7136.  
 CY United States  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 199911  
 ED Entered STN: 20000111  
 Last Updated on STN: 20000629  
 Entered Medline: 19991119  
 AB In order to analyze the **mutational** events and to understand the  
 biological significance of the p53 **gene** in chemical  
 carcinogenesis, we applied a new yeast-based p53 functional **assay**  
 to ovarian tumors induced by 7, 12-dimethylbenz[a]**anthracene**  
 (DMBA), as well as to transitional **cell** carcinomas of the  
 urinary bladder induced by N-butyl-N-(4-hydroxybutyl) nitrosamine (BBN) in  
 rats. The **assay** demonstrated that 15 of 19 DMBA induced tumors  
 harbored clonal p53 **mutations**, which is consistent with the  
 expectations of the "clonal expansion" hypothesis. The majority of the  
**mutations** were purine (AG) to pyrimidine (CT) transversions  
 (12/19) on the non-transcribed (sense) strand (NTS), which is likely to be  
 due to depurination created by DMBA adduct formation on the NTS. In  
 contrast, we found no pyrimidine to purine [corrected] transversion on the  
 NTS. After cessation of BBN treatment, BBN-induced multifocal lesions in  
 the bladder contained heterogeneous p53 **mutations** at an early  
 stage. In the later stage, however, clonal p53 **mutations** were  
 identified in 4 out of 7 bladders analyzed, conforming with the concept of  
 "field cancerization". The observed base substitutions were G-->A (1/6) or  
 C -->T transitions (2/6), and **mutations** at T (3/6) on the NTS in  
 clonal **mutations**, together with non-clonal **mutations**,  
 showing a preference of C-->T to G-->A (17 vs. 0). Thus, preferential  
 repair was found in the transcribed strand of the p53 **gene**,  
 whether modified by DMBA or by BBN carcinogens. Very similar  
**mutation** patterns were observed between clonal and non-clonal  
**mutations** in the DMBA- and BBN-induced tumors, indicating that the  
**rat** yeast p53 functional **assay** can be a potential tool  
 for the characterization of in **vivo** **mutation** patterns  
 of p53, when modified by chemical carcinogens.  
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L21 ANSWER 7 OF 56 MEDLINE  
 AN 1999453355 MEDLINE  
 DN 99453355 PubMed ID: 10521669  
 TI Induction of lacZ **mutation** by 7,12-dimethylbenz[a]  
**anthracene** in various tissues of transgenic mice.  
 AU Hachiya N; Yajima N; Hatakeyama S; Yuno K; Okada N; Umeda Y; Wakata A;  
 Motohashi Y  
 CS Department of Public Health, Akita University School of Medicine, Hondo  
 1-chome, Akita, Japan.. hachiya@ipc.akita-u.ac.jp  
 SO MUTATION RESEARCH, (1999 Aug 18) 444 (2) 283-95.  
 Journal code: 0400763. ISSN: 0027-5107.  
 CY Netherlands  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 199910  
 ED Entered STN: 20000111  
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 Entered Medline: 19991029  
 AB The induction of **gene mutations** was examined in  
**MutaMouse** after an intraperitoneal injection of 7,  
 8-dimethylbenz[a]**anthracene** (DMBA) at 20 mg/kg in a  
 collaborative study participated by four laboratories. Although the DMBA  
 dose used was lower than the level that has been reported to induce  
 micronucleated erythrocytes maximally in several **mouse** strains,  
 a killing effect appeared after day 9 of the post-treatment interval.  
**Mutations** in lacZ transgene were detected by the positive  
 selection **assay** following in vitro packaging of phage lambda  
 from the genomic DNA of the transgenic animals that survived. The  
**mutant** induction was evaluated in the bone marrow, liver, skin,  
 colon, kidney, thymus, and testis 7 to 28 days after the treatment. In the  
 bone marrow, the **mutant** frequency reached a maximum,  
 approximately a 30-fold increase, 14 days after the treatment and the  
 increased frequency persisted at least up to day 28 of the post-treatment.  
 Induction of **mutants** was detected in the liver, colon, thymus,  
 and skin to lesser extents. Marginal responses were obtained in the kidney  
 and testis. The slight increases in the **mutant** frequencies in  
 the kidney and testis observed in some laboratories were within  
 laboratory-to-laboratory or animal-to-animal variations. In contrast to  
 the **gene mutation** induction in the bone marrow, the  
 frequency of micronucleated reticulocytes increased transiently 3 days  
 after the treatment and returned to a control level before day 8 of the  
 post-treatment. It was suggested that DMBA induced **gene**  
**mutation** is fixed in stem **cells** depending on  
**cell** proliferation while DNA damages responsible for chromosome  
 breakage are not transmitted to progeny **cells**.

L21 ANSWER 6 OF 56 MEDLINE  
 AN 2000035347 MEDLINE  
 DN 20035347 PubMed ID: 10567037  
 TI Antimutagenic effects of centchroman--a contraceptive and a candidate drug for breast cancer in multiple **mutational** assays.  
 AU Giri A K; Mukhopadhyay A; Sun J; Hsie A W; Ray S  
 CS Indian Institute of Chemical Biology, 4 Raja S.C.Mullick Road, Jadavpur, Calcutta 700 032, India.. iichbio@gisclo1.vsnl.net.in  
 SO MUTAGENESIS, (1999 Nov) 14 (6) 613-20.  
 Journal code: 8707812. ISSN: 0267-8357.  
 CY ENGLAND: United Kingdom  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 200003  
 ED Entered STN: 20000320  
 Last Updated on STN: 20000320  
 Entered Medline: 20000308  
 AB Centchroman (CC), a non-steroidal oral contraceptive and a candidate drug for breast cancer, has been reported to exhibit partial to complete remission of lesions in 40.5% of breast cancer patients. The potent anti-oestrogenic activity, negligible side-effects and anti-breast cancer activity of CC prompted us to evaluate the antimutagenic effects of this compound in a bacterial **mutagenicity assay** and CHO/HPRT and AS52/GPT **mutation** assays in vitro and in **vivo** in female Swiss albino mice as measured by both sister chromatid exchange (SCE) and chromosome aberrations (CA) against three known positive **mutagen** compounds, dimethylbenz[a]**anthracene** (DMBA), cyclophosphamide (CP) and mitomycin C (MMC). Antimutagenicity assays in Salmonella strains TA97a, TA100, TA98 and TA102 were carried out against commonly used known positive **mutagens**, sodium azide, 4-nitro-o-phenylenediamine, cumene hydroperoxide, 2-aminofluorene and danthron. A significantly reduced number of bacterial histidine revertant colonies was observed in the plates treated with 0.1, 1, 5 and 10 microg/plate CC and a positive compound when compared with bacterial plates treated with the respective positive compound alone. Ethyl methanesulfonate (EMS), a commonly used positive **mutagen** for CHO/HPRT and AS52/GPT **gene mutation** assays, was used for antimutagenicity **assay** in these **cells**. CC exhibited protective effects against the **mutagenicity** of EMS in these two mammalian **cell mutation** assays, CHO/HPRT and AS52/GPT. In the in **vivo** studies, pretreatment with CC reduced DMBA-induced SCE and CA and CP- and MMC-induced CA when compared with the group treated only with the positive compounds. These results indicate that CC can reduce the **mutagenic** effects of known genotoxic compounds.

L21 ANSWER 5 OF 56 MEDLINE  
 AN 2000299169 MEDLINE  
 DN 20299169 PubMed ID: 10838135  
 TI **Mutational** spectra for polycyclic aromatic hydrocarbons in the supF target **gene**.  
 AU Bigger C A; Ponten I; Page J E; Dipple A  
 CS Chemistry of Carcinogenesis Laboratory, Basic Research Program, Advanced BioScience Laboratories, Frederick Cancer Research and Development Center, National Cancer Institute, Frederick, MD 21702, USA.. biggera@cder.fda.gov  
 SO MUTATION RESEARCH, (2000 May 30) 450 (1-2) 75-93. Ref: 77  
 Journal code: 0400763. ISSN: 0027-5107.  
 CY Netherlands  
 DT Journal; Article; (JOURNAL ARTICLE)  
 General Review; (REVIEW)  
 (REVIEW, TUTORIAL)  
 LA English  
 FS Priority Journals  
 EM 200007  
 ED Entered STN: 20000810  
 Last Updated on STN: 20000810  
 Entered Medline: 20000727  
 AB An SV40-based shuttle vector system was used to identify the types of **mutational** changes and the sites of **mutation** within the supF DNA sequence generated by the four stereoisomers of benzo[c]phenanthrene 3,4-dihydrodiol 1,2-epoxide (B[c]PhDE), by racemic mixtures of bay or fjord region dihydrodiol epoxides (DE) of 5-methylchrysene, of 5, 6-dimethylchrysene, of benzo[g]chrysene and of 7-methylbenz[a]**anthracene** and by two direct acting polycyclic aromatic hydrocarbon carcinogens, 7-bromomethylbenz[a]**anthracene** (7-BrMeBA) and 7-bromomethyl-12-methylbenz[a]**anthracene** (7-BrMe-12-MeBA). The results of these studies demonstrated that the predominant type of **mutation** induced by these compounds is the base substitution. The chemical preference for reaction at deoxyadenosine (dAdo) or deoxyguanosine (dGuo) residues in DNA, which is in general correlated with the spatial structure (planar or non-planar) of the reactive polycyclic aromatic hydrocarbon, is reflected in the preference for **mutation** at A&z.ccirf;T or G&z.ccirf;C pairs. In addition, if the ability to react with DNA in **vivo** is taken into account, the relative **mutagenic** potencies of the B[c]PhDE stereoisomers are consistent with the higher tumorigenic activity associated with non-planar polycyclic aromatic hydrocarbons and their extensive reaction with dAdo residues in DNA. Comparison of the types of **mutations** generated by polycyclic aromatic hydrocarbons and other bulky carcinogens in this shuttle vector system suggests that all bulky lesions may be processed by a similar mechanism related to that involved in replication past apurinic sites. However, inspection of the distribution of **mutations** over the target **gene** induced by the different compounds demonstrated that individual polycyclic aromatic hydrocarbons induce unique patterns of **mutational** hotspots within the target **gene**. A polymerase arrest **assay** was used to determine the sequence specificity of the interaction of reactive polycyclic aromatic hydrocarbons with the shuttle vector DNA. The results of these assays revealed a divergence between **mutational** hotspots and polymerase arrest sites for all compounds investigated, i.e., sites of **mutational** hotspots do not correspond to sites where high levels of adduct formation occur, and suggested that some association between specific adducts and sequence context may be required to constitute a premutagenic lesion. A site-specific **mutagenesis** system employing a single-stranded vector (M13mp7L2) was used to investigate the **mutational** events a single benzo[a]pyrene or benzo[c]phenanthrene dihydrodiol epoxide-DNA adduct elicits within specific sequence contexts.

These studies showed that sequence context can cause striking differences in **mutagenic** frequencies for given adducts. In addition, these sequence context effects do not originate only from nucleotides immediately adjacent to the adduct, but are also modulated by more distal nucleotides. The implications of these results for mechanisms of polycyclic aromatic hydrocarbon-induced **mutagenesis** and carcinogenesis are discussed.

L21 ANSWER 4 OF 56 MEDLINE  
 AN 2001061127 MEDLINE  
 DN 20500999 PubMed ID: 11044905  
 TI **Mutant** frequency and molecular analysis of **in vivo**  
 lacI **mutations** in the bone marrow of Big Blue rats treated with  
 7, 12-dimethylbenz[a]**anthracene**.  
 AU Shelton S D; Cherry V; Manjanatha M G  
 CS Division of Genetic Toxicology, National Center for Toxicological  
 Research, Food and Drug Administration, Jefferson, Arkansas 72079, USA.  
 SO ENVIRONMENTAL AND MOLECULAR MUTAGENESIS, (2000) 36 (3) 235-42.  
 Journal code: 8800109. ISSN: 0893-6692.  
 CY United States  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 200012  
 ED Entered STN: 20010322  
 Last Updated on STN: 20010322  
 Entered Medline: 20001222  
 AB Recently, we evaluated lacI **mutations** in lymphocytes and mammary  
 tissue of Big Blue (BB) rats exposed to 7, 12-dimethylbenz[a]  
**anthracene** (DMBA). The results on the time course of  
**mutant** induction suggested that the lacI **gene** may  
 manifest a tissue-specific increase in **mutant** frequency (MF). To  
 test whether a tissue-specific increase in lacI MF is dependent on the  
**cell** proliferation rate of a tissue, we examined rapidly  
 proliferating bone marrow **cells** for DMBA-induced lacI  
**mutations**. Seven-week-old female BB rats were given single doses  
 of 0, 20, and 130 mg/kg DMBA by gavage and the lacI MFs in the bone marrow  
 were measured over a period of 14 weeks following treatment. Bone marrow  
**cells** had a remarkably low average background MF (3.1 +/- 1.6 x  
 10<sup>-6</sup> plaque-forming units) and the DMBA-induced lacI MFs were  
 significantly higher than control MFs for both doses and at all time  
 points (P < 0.01). The lacI MF in the bone marrow increased for 2 weeks  
 and then remained relatively constant; 20 and 130 mg/kg DMBA produced 34-  
 and 106-fold increases in MF over control MF. DNA sequencing revealed that  
 the majority of DMBA-induced lacI **mutations** were base-pair  
 substitutions and that A:T --> T:A (48%) and G:C --> T:A (24%)  
 transversions were the predominant types. Thus, the different lacI  
**mutation** fixation times observed for bone marrow (2 weeks),  
 mammary (10 weeks), and lymphocytes (6 weeks) suggest that the lacI  
**gene** manifests a tissue-specific **mutation** fixation time,  
 which may depend on the **cell** proliferation rate of a tissue. In  
 addition, the relatively low spontaneous MF in bone marrow compared with  
 that in other tissues may be useful for increasing the sensitivity of the  
**assay** for detecting induced MFs in BB rats.

L21 ANSWER 3 OF 56 MEDLINE  
 AN 2001098636 MEDLINE  
 DN 20584309 PubMed ID: 11152561  
 TI 7,12-dimethylbenz[a]anthracene-induced **mutation** in the  
 Tk **gene** of Tk(+/-) mice: automated scoring of lymphocyte clones  
 using a fluorescent viability indicator.  
 AU Dobrovolsky V N; Shaddock J G; Heflich R H  
 CS Division of Genetic and Reproductive Toxicology, HFT-120, National Center  
 for Toxicological Research, Jefferson, Arkansas 72079, USA..  
 vdobrovolsky@nctr.fda.gov  
 SO ENVIRONMENTAL AND MOLECULAR MUTAGENESIS, (2000) 36 (4) 283-91.  
 Journal code: 8800109. ISSN: 0893-6692.  
 CY United States  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 200102  
 ED Entered STN: 20010322  
 Last Updated on STN: 20010322  
 Entered Medline: 20010201  
 AB 7,12-Dimethylbenz[a]anthracene (DMBA) is a rodent carcinogen and  
 a potent **in vivo mutagen** for the X-linked hypoxanthine  
 guanine phosphoribosyl transferase (hprt) **gene** of rats and for  
 the lacI transgene of Big Blue mice and rats. Although DMBA is also a  
 powerful clastogen, molecular analysis of these DMBA-induced hprt and lacI  
**mutations** indicates that most are single base-pair (bp)  
 substitutions and 1- to 3-bp frameshifts. In the present study, we  
 evaluated the types of **mutations** induced by DMBA in the  
 autosomal thymidine kinase (Tk) **gene** of Tk(+/-) mice. Male and  
 female 5- to 6-week-old animals were injected i.p. with DMBA at a dose of  
 30 mg/kg. Five weeks after the treatment, hprt and Tk **mutant**  
 frequencies were determined using a limiting dilution clonal **assay**  
 in 96-well plates. We established conditions for the automated  
 identification of wells containing expanded lymphocyte clones using the  
 fluorescent indicator alamarBlue. This procedure allowed the unbiased  
 identification of viable clones and calculation of **mutant**  
 frequencies. In male mice, DMBA treatment increased the frequency of hprt  
**mutants** from 1.8 +/- 1.1 to 34 +/- 9 x 10<sup>(-6)</sup>, and Tk  
**mutants** from 33 +/- 12 to 78 +/- 26 x 10<sup>(-6)</sup>; treated female mice  
 had a significant but lower increase in hprt **mutant** frequency  
 than did males. Molecular analysis of DMBA-induced Tk **mutants**  
 revealed that at least 75% had the entire wild-type Tk allele missing. The  
 results indicate that the predominant types of DMBA-induced  
**mutation** detected by the autosomal Tk **gene** are different  
 from those detected by the X-linked hprt **gene**. The Tk  
**gene** mainly detects loss of heterozygosity **mutation**,  
 whereas the majority of **mutations** previously found in the hprt  
**gene** were point **mutations**.